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764

1

SEQUENCE LISTING

<110> OSTERMEIER, MARK

<120> MOLECULAR SWITCHES AND METHODS FOR MAKING AND USING THE  
SAME

<130> 56908 (71699)

<140> 10/507,466

<141> 2004-09-10

<150> PCT/US03/07380

<151> 2003-03-10

<150> 60/362,588

<151> 2002-03-11

<160> 41

<170> PatentIn Ver. 3.3

<210> 1

<211> 4

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic  
linker sequence

<400> 1

Gly Ser Gly Gly

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<210> 2

<211> 37

<212> PRT

<213> Homo sapiens

<400> 2

Pro Asn Lys Gly Ser Gly Thr Thr Ser Gly Thr Thr Arg Leu Leu Ser

1

5

10

15

Gly His Thr Cys Phe Thr Leu Thr Gly Leu Leu Gly Thr Leu Val Thr

20

25

30

Met Gly Leu Leu Thr

35

<210> 3

<211> 14

<212> PRT

<213> Unknown Organism

<220>

<223> Description of Unknown Organism: Myristylation  
src amino acid sequence

<400> 3

Met Gly Ser Ser Lys Ser Lys Pro Lys Asp Pro Ser Gln Arg  
1 5 10

<210> 4

<211> 25

<212> PRT

<213> Unknown Organism

<220>

<223> Description of Unknown Organism: Palmitoylation GRK6  
amino acid sequence

<400> 4

Leu Leu Gln Arg Leu Phe Ser Arg Gln Asp Cys Cys Gly Asn Cys Ser  
1 5 10 15

Asp Ser Glu Glu Glu Leu Pro Thr Arg  
20 25

<210> 5

<211> 7

<212> PRT

<213> Monkey virus

<400> 5

Pro Lys Lys Lys Lys Val  
1 5

<210> 6

<211> 6

<212> PRT

<213> Homo sapiens

<400> 6

Ala Arg Arg Arg Arg Pro  
1 5

<210> 7

<211> 10

<212> PRT

<213> Unknown Organism

<220>

<223> Description of Unknown Organism: NF kB p50 amino  
acid sequence

<400> 7

Glu Glu Val Gln Arg Lys Arg Gln Lys Leu  
1 5 10

<210> 8  
 <211> 9  
 <212> PRT  
 <213> Unknown Organism

<220>  
 <223> Description of Unknown Organism: NF kB p65 amino  
 acid sequence

<400> 8  
 Glu Glu Lys Arg Lys Arg Thr Tyr Glu  
 1 5

<210> 9  
 <211> 21  
 <212> PRT  
 <213> Unknown Organism

<220>  
 <223> Description of Unknown Organism: Nucleoplasmin  
 amino acid sequence

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 Ala Val Lys Arg Pro Ala Ala Thr Leu Lys Lys Ala Gly Gln Ala Lys  
 1 5 10 15

Lys Lys Lys Leu Asp  
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<210> 10  
 <211> 5  
 <212> PRT  
 <213> Homo sapiens

<400> 10  
 Lys Phe Glu Arg Gln  
 1 5

<210> 11  
 <211> 36  
 <212> PRT  
 <213> Homo sapiens

<400> 11  
 Met Leu Ile Pro Ile Ala Gly Phe Phe Ala Leu Ala Gly Leu Val Leu  
 1 5 10 15

Ile Val Leu Ile Ala Tyr Leu Ile Gly Arg Lys Arg Ser His Ala Gly  
 20 25 30

Tyr Gln Thr Ile  
 35

<210> 12  
 <211> 35  
 <212> PRT  
 <213> Homo sapiens

<400> 12  
 Leu Val Pro Ile Ala Val Gly Ala Ala Leu Ala Gly Val Leu Ile Leu  
   1                  5                  10                  15  
 Val Leu Leu Ala Tyr Phe Ile Gly Leu Lys His His His Ala Gly Tyr  
                   20                  25                  30  
 Glu Gln Phe  
                   35

<210> 13  
 <211> 27  
 <212> PRT  
 <213> Saccharomyces sp.

<400> 13  
 Met Leu Arg Thr Ser Ser Leu Phe Thr Arg Arg Val Gln Pro Ser Leu  
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 Phe Ser Arg Asn Ile Leu Arg Leu Gln Ser Thr  
                   20                  25

<210> 14  
 <211> 25  
 <212> PRT  
 <213> Saccharomyces sp.

<400> 14  
 Met Leu Ser Leu Arg Gln Ser Ile Arg Phe Phe Lys Pro Ala Thr Arg  
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 Thr Leu Cys Ser Ser Arg Tyr Leu Leu  
                   20                  25

<210> 15  
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 <212> PRT  
 <213> Saccharomyces sp.

<400> 15  
 Met Phe Ser Met Leu Ser Lys Arg Trp Ala Gln Arg Thr Leu Ser Lys  
   1                  5                  10                  15  
 Ser Phe Tyr Ser Thr Ala Thr Gly Ala Ala Ser Lys Ser Gly Lys Leu  
                   20                  25                  30  
 Thr Gln Lys Leu Val Thr Ala Gly Val Ala Ala Ala Gly Ile Thr Ala  
                   35                  40                  45

Ser Thr Leu Leu Tyr Ala Asp Ser Leu Thr Ala Glu Ala Met Thr Ala  
 50 55 60

<210> 16  
 <211> 41  
 <212> PRT  
 <213> Saccharomyces sp.

<400> 16  
 Met Lys Ser Phe Ile Thr Arg Asn Lys Thr Ala Ile Leu Ala Thr Val  
 1 5 10 15  
 Ala Ala Thr Gly Thr Ala Ile Gly Ala Tyr Tyr Tyr Tyr Asn Gln Leu  
 20 25 30  
 Gln Gln Gln Gln Gln Arg Gly Lys Lys  
 35 40

<210> 17  
 <211> 4  
 <212> PRT  
 <213> Unknown Organism

<220>  
 <223> Description of Unknown Organism: Protein sequence  
 from calreticulin

<400> 17  
 Lys Asp Glu Leu  
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<210> 18  
 <211> 15  
 <212> PRT  
 <213> Human adenovirus

<400> 18  
 Leu Tyr Leu Ser Arg Arg Ser Phe Ile Asp Glu Lys Lys Met Pro  
 1 5 10 15

<210> 19  
 <211> 20  
 <212> PRT  
 <213> Homo sapiens

<400> 19  
 Met Tyr Arg Met Gln Leu Leu Ser Cys Ile Ala Leu Ser Leu Ala Leu  
 1 5 10 15  
 Val Thr Asn Ser  
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<210> 20  
 <211> 29  
 <212> PRT  
 <213> Homo sapiens

<400> 20  
 Met Ala Thr Gly Ser Arg Thr Ser Leu Leu Leu Ala Phe Gly Leu Leu  
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 Cys Leu Pro Trp Leu Gln Glu Gly Ser Ala Phe Pro Thr  
                           20                          25

<210> 21  
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 <212> PRT  
 <213> Homo sapiens

<400> 21  
 Met Ala Leu Trp Met Arg Leu Leu Pro Leu Leu Ala Leu Leu Ala Leu  
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 Trp Gly Pro Asp Pro Ala Ala Ala Phe Val Asn  
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<210> 22  
 <211> 18  
 <212> PRT  
 <213> Influenza virus

<400> 22  
 Met Lys Ala Lys Leu Leu Val Leu Leu Tyr Ala Phe Val Ala Gly Asp  
           1                          5                          10                          15  
 Gln Ile

<210> 23  
 <211> 24  
 <212> PRT  
 <213> Homo sapiens

<400> 23  
 Met Gly Leu Thr Ser Gln Leu Leu Pro Pro Leu Phe Phe Leu Leu Ala  
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 Cys Ala Gly Asn Phe Val His Gly  
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<210> 24  
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 <212> PRT  
 <213> Artificial Sequence

<220>  
 <223> Description of Artificial Sequence: Synthetic  
 linker

<400> 24  
 Gly Gly Gly Ser  
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<210> 25  
 <211> 4  
 <212> PRT  
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<220>  
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<400> 25  
 Ser Gly Gly Gly  
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<210> 26  
 <211> 12  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Description of Artificial Sequence: Synthetic  
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<400> 26  
 ggtggtggca gc

12

<210> 27  
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<220>  
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<400> 27  
 agcgtggcg gc

12

<210> 28  
 <211> 8  
 <212> PRT  
 <213> Artificial Sequence

<220>  
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 linker

<400> 28

Gly Ser Gly Gly Gly Ser Gly Gly  
1 5

<210> 29

<211> 35

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic  
oligonucleotide

<400> 29

gccttcagat ctcttctcac ccagaaacgc tgggtg

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<210> 30

<211> 35

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic  
oligonucleotide

<400> 30

tcactgatta agcattggtg aagagatctg gttca

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<210> 31

<211> 35

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic  
oligonucleotide

<400> 31

caccagcggtt tctgggtgag aagagatctg aaggc

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<210> 32

<211> 35

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic  
oligonucleotide

<400> 32

tgaaccagat ctcttcactt ggtgatacga gtctg

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<210> 33  
 <211> 18  
 <212> DNA  
 <213> Artificial Sequence

<220>  
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<400> 33  
 caccagaaa cgctggtg 18

<210> 34  
 <211> 15  
 <212> DNA  
 <213> Artificial Sequence

<220>  
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<400> 34  
 tcactgatta agcat 15

<210> 35  
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<220>  
 <223> Description of Artificial Sequence: Synthetic  
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<400> 35  
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<210> 36  
 <211> 18  
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<220>  
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<400> 36  
 cttggtgata cgagtctg 18

<210> 37  
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<220>  
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oligonucleotide

<400> 37  
tcactgatta agcataag 18

<210> 38  
<211> 18  
<212> DNA  
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<220>  
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oligonucleotide

<400> 38  
caccagcgtt tctgggtg 18

<210> 39  
<211> 7  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: Synthetic  
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<400> 39  
ctgatcc 7

<210> 40  
<211> 10  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: Synthetic  
oligonucleotide

<400> 40  
gagacggcga 10

<210> 41  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: Synthetic  
oligonucleotide construct

<400> 41  
ctgatcgcta ggagacggtg c 21